

Refine Search

Your wildcard search against 10000 terms has yielded the results below.

Your result set for the last L# is incomplete.

The probable cause is use of unlimited truncation. Revise your search strategy to use limited truncation.

Search Results -

Terms	Documents
enhance near export near5 (mRNA\$ or RNA\$) near20 (retrovir\$ or adenovir\$ or lentivir\$ or herpesvir\$ or viral or virus or viruses)	2

Database:

US Pre-Grant Publication Full-Text Database
US Patents Full-Text Database
US OCR Full-Text Database
EPO Abstracts Database
JPO Abstracts Database
Derwent World Patents Index
IBM Technical Disclosure Bulletins

Search:

L10

Refine Search

Recall Text

Clear

Interrupt

Search History

DATE: Friday, October 27, 2006 [Purge Queries](#) [Printable Copy](#) [Create Case](#)

<u>Set Name</u> side by side	<u>Query</u>	<u>Hit Count</u>	<u>Set Name</u> result set
<i>DB=PGPB,USPT,USOC,EPAB,JPAB,DWPI,TDBD; PLUR=YES; OP=OR</i>			
<u>L10</u>	enhance near export near5 (mRNA\$ or RNA\$) near20 (retrovir\$ or adenovir\$ or lentivir\$ or herpesvir\$ or viral or virus or viruses)	2	<u>L10</u>
<u>L9</u>	enhance near export near5 (mRNA\$ or RNA\$) near20 (retrovir\$ or adenovir\$ or lentivir\$ or herpesvir\$)	0	<u>L9</u>
<u>L8</u>	mRNA near processing near enhancer	24	<u>L8</u>
<u>L7</u>	PPE near20 (retrovir\$ or adenovir\$ or lentivir\$ or herpesvir\$)	8	<u>L7</u>
<u>L6</u>	mRNA near5 processing near5 enhancer near50 (retrovir\$ or adenovir\$ or lentivir\$ or herpesvir\$)	1	<u>L6</u>
<u>L5</u>	mRNA near5 processing near5 enhancer	56	<u>L5</u>
<u>L4</u>	tat near10 (fused or fusion\$ or substitut\$) near10 HTLV-1	0	<u>L4</u>

L3 tat near10 (fused or fusion\$ or substitut\$)
L2 tat near10 (fused or fusion\$ or substitut\$) near10 Tax
L1 tat near10 (fused or fusion\$ or substitut\$) Tax

1211 L3
1 L2
28068 L1

END OF SEARCH HISTORY



Day : Friday
Date: 10/27/2006

Time: 15:31:26

Inventor Name Search

Enter the **first few letters** of the Inventor's Last Name.
Additionally, enter the **first few letters** of the Inventor's First name.

Last Name

First Name

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Inventor Name Search

Enter the **first few letters** of the Inventor's Last Name.
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Additionally, enter the **first few letters** of the Inventor's First name.

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**PALM INTRANET**Day : Friday
Date: 10/27/2006

Time: 15:31:26

Inventor Name Search

Enter the **first few letters** of the Inventor's Last Name.
Additionally, enter the **first few letters** of the Inventor's First name.

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Search	Most Recent Queries	Time	Result
#12	Search inducible retroviral promoter and tax and tat and wpre	15:59:27	0
#11	Search inducible retroviral promoter and tax and tat	15:58:16	1
#10	Search inducible retroviral promoter and tax	15:58:11	39
#9	Search tat tax chimera	15:57:08	1
#8	Search tat tax hybrid	15:56:39	6
#1	Search tat fusion tax	15:54:32	8

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Oct 24 2006 07:33:51

Set	Items	Description
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? set hi ;set hi

HIGHLIGHT set on as ''

HIGHLIGHT set on as ''

? begin 5,6,55,154,155,156,312,399,biotech,biosci

>>> 44 is unauthorized

Set Items Description

? s induc? (5n) retrovir? (5n) promoter?

Processing

Processed 10 of 41 files ...

Processing

Completed processing all files

14170972 INDUC?

697262 RETROVIR?

1321776 PROMOTER?

S1 296 INDUC? (5N) RETROVIR? (5N) PROMOTER?

? s s1 and tax

296 S1

90442 TAX

S2 6 S1 AND TAX

? s s2 and tat

6 S2

72043 TAT

S3 3 S2 AND TAT

? s s1 and (WPRE or woodchuck)

296 S1

574 WPRE

7826 WOODCHUCK

S4 2 S1 AND (WPRE OR WOODCHUCK)

? s s4/3/1-2

>>>Invalid syntax

? d s4/3/1-2

Display 4/3/1 (Item 1 from file: 399)

DIALOG(R)File 399:CA SEARCH(R)

(c) 2006 American Chemical Society. All rts. reserv.

142192342 CA: 142(11)192342h PATENT

Inducible mammalian protein expression system comprising a retroviral promoter and a promoter-activating protein

INVENTOR(AUTHOR): Harms, Jerome S.; Splitter, Gary A.; Eakle, Kurt A.; Bremel, Robert D.

LOCATION: USA

PATENT: U.S. Pat. Appl. Publ. ; US 20050026288 A1 DATE: 20050203

APPLICATION: US 2004763976 (20040123) *US 2003PV442103 (20030123)

PAGES: 117 pp. CODEN: USXXCO LANGUAGE: English

PATENT CLASSIFICATIONS:

CLASS: 435456000; C12N-015/86A

- end of record -

?

Display 4/3/2 (Item 1 from file: 357)

DIALOG(R)File 357:Derwent Biotech Res.

(c) 2006 The Thomson Corp. All rts. reserv.

0362159 DBR Accession No.: 2005-07863 PATENT

New inducible gene expression system comprising a first vector comprising at least one retroviral promoter, at least one factor inducing the retroviral promoter, and at least one gene product, useful for expressing genes and proteins - plasmid, cosmid or virus vector-mediated retro virus promoter gene transfer and expression in host cell for protein and gene expression and gene therapy

AUTHOR: HARMS J S; SPLITTER G A; EAKLE K A; BREMEL R D

PATENT ASSIGNEE: HARMS J S; SPLITTER G A; EAKLE K A; BREMEL R D 2005

PATENT NUMBER: US 20050026288 PATENT DATE: 20050203 WPI ACCESSION NO.:

2005-141386 (200515)

PRIORITY APPLIC. NO.: US 763976 APPLIC. DATE: 20040123

NATIONAL APPLIC. NO.: US 763976 APPLIC. DATE: 20040123

LANGUAGE: English

- end of record -

? s Tax (5n) (fused or fusion or fusions) (5n) tat

90442 TAX
375242 FUSED
1350350 FUSION
80283 FUSIONS
72043 TAT

S5 13 TAX (5N) (FUSED OR FUSION OR FUSIONS) (5N) TAT

? rd s5

>>>Duplicate detection is not supported for File 391.

>>>Records from unsupported files will be retained in the RD set.

S6 4 RD S5 (unique items)

? d s6/3/1-4

Display 6/3/1 (Item 1 from file: 5)

DIALOG(R)File 5:Biosis Previews(R)

(c) 2006 The Thomson Corporation. All rts. reserv.

0012499960 BIOSIS NO.: 200000218273

Human T-cell leukemia virus type 1 Tax shuttles between functionally
discrete subcellular targets

AUTHOR: Burton Molly; Upadhyaya Cherrag D; Maier Bernhard; Hope Thomas J;
Semmes O John (Reprint)

AUTHOR ADDRESS: Department of Microbiology, University of Virginia School
of Medicine, Jordan Hall 7-89, Charlottesville, VA, 23060, USA**USA

JOURNAL: Journal of Virology 74 (5): p2351-2364 March, 2000 2000

MEDIUM: print

ISSN: 0022-538X

DOCUMENT TYPE: Article

RECORD TYPE: Abstract

LANGUAGE: English

- end of record -

?

Display 6/3/2 (Item 1 from file: 154)

DIALOG(R)File 154:MEDLINE(R)

(c) format only 2006 Dialog. All rts. reserv.

10597877 PMID: 7474143

Selective infection of human T-lymphotropic virus type 1
(HTLV-1)-infected cells by chimeric human immunodeficiency viruses
containing HTLV-1 tax response elements in the long terminal repeat:

Lin H C; Bodkin M; Lal R B; Rabson A B

Department of Molecular Genetics and Microbiology, Robert Wood Johnson
Medical School, University of Medicine and Dentistry of New Jersey,
Piscataway, USA.

Journal of virology (UNITED STATES) Nov 1995, 69 (11) p7216-25,

ISSN 0022-538X--Print Journal Code: 0113724

Contract/Grant No.: AI30901; AI; NIAID

Publishing Model Print

Document type: Journal Article

Languages: ENGLISH

Main Citation Owner: NLM

Record type: MEDLINE; Completed

- end of record -

?

Display 6/3/3 (Item 1 from file: 399)

DIALOG(R)File 399:CA SEARCH(R)

(c) 2006 American Chemical Society. All rts. reserv.

142192342 CA: 142(11)192342h PATENT
Inducible mammalian protein expression system comprising a retroviral
promoter and a promoter-activating protein
INVENTOR(AUTHOR): Harms, Jerome S.; Splitter, Gary A.; Eakle, Kurt A.;
Bremel, Robert D.
LOCATION: USA
PATENT: U.S. Pat. Appl. Publ. ; US 20050026288 A1 DATE: 20050203
APPLICATION: US 2004763976 (20040123) *US 2003PV442103 (20030123)
PAGES: 117 pp. CODEN: USXXCO LANGUAGE: English
PATENT CLASSIFICATIONS:
CLASS: 435456000; C12N-015/86A

- end of record -

?
Display 6/3/4 (Item 1 from file: 357)
DIALOG(R)File 357:Derwent Biotech Res.
(c) 2006 The Thomson Corp. All rts. reserv.

0362159 DBR Accession No.: 2005-07863 PATENT
New inducible gene expression system comprising a first vector comprising
at least one retroviral promoter, at least one factor inducing the
retroviral promoter, and at least one gene product, useful for
expressing genes and proteins - plasmid, cosmid or virus
vector-mediated retro virus promoter gene transfer and expression in
host cell for protein and gene expression and gene therapy
AUTHOR: HARMS J S; SPLITTER G A; EAKLE K A; BREMEL R D
PATENT ASSIGNEE: HARMS J S; SPLITTER G A; EAKLE K A; BREMEL R D 2005
PATENT NUMBER: US 20050026288 PATENT DATE: 20050203 WPI ACCESSION NO.:
2005-141386 (200515)
PRIORITY APPLIC. NO.: US 763976 APPLIC. DATE: 20040123
NATIONAL APPLIC. NO.: US 763976 APPLIC. DATE: 20040123
LANGUAGE: English

- end of record -

?
? d s6/9/1-2
Display 6/9/1 (Item 1 from file: 5)
DIALOG(R)File 5:Biosis Previews(R)
(c) 2006 The Thomson Corporation. All rts. reserv.

0012499960 BIOSIS NO.: 200000218273
Human T-cell leukemia virus type 1 Tax shuttles between functionally
discrete subcellular targets
AUTHOR: Burton Molly; Upadhyaya Cherrag D; Maier Bernhard; Hope Thomas J;
Semmes O John (Reprint)
AUTHOR ADDRESS: Department of Microbiology, University of Virginia School
of Medicine, Jordan Hall 7-89, Charlottesville, VA, 23060, USA**USA
JOURNAL: Journal of Virology 74 (5): p2351-2364 March, 2000 2000
MEDIUM: print
ISSN: 0022-538X
DOCUMENT TYPE: Article
RECORD TYPE: Abstract
LANGUAGE: English

ABSTRACT: Human T-cell leukemia virus type 1 (HTLV-1) Tax is a nuclear
protein with striking pleiotropic functionality. We recently demonstrated

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Display 6/9/1 (Item 1 from file: 5)
DIALOG(R)File 5:Biosis Previews(R)
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that Tax localizes to a multicomponent nuclear speckled structure (Tax
speckled structure (TSS)). Here, we examine these structures further and

identify a partial overlap of TSS with transcription hot spots. We used a strategy of directed expression via fusion proteins to determine if these transcription sites are the subtargets within TSS required for Tax function. When ***fused*** to human immunodeficiency virus type 1 (HIV-1) Tat, the resulting Tat-Tax fusion protein displayed neither a Tat-like nor a Tax-like pattern but rather was targeted specifically to the transcription subsites. The Tat-Tax fusion was able to activate both the HIV-1 long terminal repeat (LTR) and the HTVL-1 LTR at the same level as the individual component; thus, targeting proteins to transcription hot spots was compatible with both ***Tax*** and ***Tat*** transcription function. In contrast, the fusion with HIV-1 Rev, Rev-Tax, resulted in a pattern of expression that was largely Rev-like (nucleolar and cytoplasmic). The reduced localization of Rev-Tax to transcription sites was reflected in a 10-fold drop in activation of the HTLV-1 LTR. However,

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DIALOG(R)File 5:Biosis Previews(R)
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there was no loss in the ability of Tax to activate via NF-kappaB. Thus, NF-kappaB-dependent Tax function does not require targeting of Tax to these transcription sites and suggests that activation via NF-kappaB is a cytoplasmic function. Selective mutation of the nuclear localization signal site in the Rev portion resulted in retargeting of Rev-Tax to TSS and subsequent restoration of transcription function, demonstrating that inappropriate localization preceded loss of function. Mutation of the nuclear export signal site in the Rev portion had no effect on transcription, although the relative amount of Rev-Tax in the cytoplasm was reduced. Finally, in explaining how Tax can occupy multiple subcellular sites, we show that Tax shuttles from the nucleus to the cytoplasm in a heterokaryon fusion assay. Thus, pleiotropic functionality by Tax is regulatable via shuttling between discrete subcellular compartments.

DESCRIPTORS:

MAJOR CONCEPTS: Molecular Genetics--Biochemistry and Molecular Biophysics

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Display 6/9/1 (Item 1 from file: 5)
DIALOG(R)File 5:Biosis Previews(R)
(c) 2006 The Thomson Corporation. All rts. reserv.
; Infection
BIOSYSTEMATIC NAMES: Hominidae--Primates, Mammalia, Vertebrata, Chordata, Animalia; Retroviridae--DNA and RNA Reverse Transcribing Viruses, Viruses, Microorganisms
ORGANISMS: HeLa cell line (Hominidae); human T-cell leukemia virus type 1 (Retroviridae)
ORGANISMS: PARTS ETC: cytoplasm; nucleolus
COMMON TAXONOMIC TERMS: Animals; Chordates; Humans; Mammals; Primates; Vertebrates; DNA and RNA Reverse Transcribing Viruses; Microorganisms; Viruses
CHEMICALS & BIOCHEMICALS: Tax proteins; fusion proteins
MISCELLANEOUS TERMS: pleiotropy; subcellular compartments; transcription; transcription hot spots; transcription subsites
CONCEPT CODES:
33506 Virology - Animal host viruses
02508 Cytology - Human
10064 Biochemistry studies - Proteins, peptides and amino acids

-more-

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Display 6/9/1 (Item 1 from file: 5)
DIALOG(R)File 5:Biosis Previews(R)
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10506 Biophysics - Molecular properties and macromolecules
10508 Biophysics - Membrane phenomena
11108 Anatomy and Histology - Microscopic and ultramicroscopic anatomy
36006 Medical and clinical microbiology - Virology
12502 Pathology - General
13012 Metabolism - Proteins, peptides and amino acids
31500 Genetics of bacteria and viruses
32000 Microbiological apparatus, methods and media
BIOSYSTEMATIC CODES:
86215 Hominidae
03305 Retroviridae

- end of record -

?

Display 6/9/2 (Item 1 from file: 154)
DIALOG(R)File 154:MEDLINE(R)
(c) format only 2006 Dialog. All rts. reserv.
10597877 PMID: 7474143
Selective infection of human T-lymphotropic virus type 1 (HTLV-1)-infected cells by chimeric human immunodeficiency viruses containing HTLV-1 tax response elements in the long terminal repeat.
Lin H C; Bodkin M; Lal R B; Rabson A B
Department of Molecular Genetics and Microbiology, Robert Wood Johnson Medical School, University of Medicine and Dentistry of New Jersey, Piscataway, USA.
Journal of virology (UNITED STATES) Nov 1995, 69 (11) p7216-25,
ISSN 0022-538X--Print Journal Code: 0113724
Contract/Grant Number: AI30901; AI; NIAID
Publishing Model Print
Document type: Journal Article
Languages: ENGLISH
Main Citation Owner: NLM
Record type: MEDLINE; Completed

-more-

?

Display 6/9/2 (Item 1 from file: 154)
DIALOG(R)File 154:MEDLINE(R)
(c) format only 2006 Dialog. All rts. reserv.
Subfile: INDEX MEDICUS; AIDS/HIV; Toxbib
Previous studies have suggested that the human immunodeficiency virus long terminal repeat (HIV LTR) enhancer/promoter sequences contribute to the replication ability of HIV in different T-cell lines; mutation of these sequences can alter HIV tropism. We have utilized site-specific mutagenesis to generate variants of HIV that exhibit specific tropism for human T-lymphotropic virus type 1 (HTLV-1) Tax-expressing CD4+ T cells. The wild-type HIV LTR NF-kappa B and Spl sites in an infectious molecular clone of HIV type 1 were replaced with sequences derived from the 21-bp Tax response elements (TRE) from the HTLV-1 LTR to generate TRE-containing chimeric HIVs (TRE-HIVs). The TRE-HIVs exhibit selective replication and cell killing in HTLV-infected human CD4+ T cells, but not in HTLV-negative T cells. Transient transfections suggested that Tax-TRE interactions could account for the observed replication specificity. The TRE-containing HIV LTRs were synergistically activated by the HIV Tat and HTLV-1 Tax transactivators. These results demonstrate that it is possible to specifically target HIV replication and cytotoxicity to HTLV-1+, CD4+ human

-more-

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DIALOG(R) File 154:MEDLINE(R)

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T cells, on the basis of Tax-TRE interactions, and provide a model for the development of specific, cytotoxic, retroviral gene therapy vectors for HTLV-1-infected cells based on alterations of the LTR transcriptional regulatory elements. They also suggest that HIV Tat can cooperate with heterologous transcriptional activators, such as Tax, which act through upstream binding sites without directly binding to DNA.

Descriptors: *Gene Products, tat--metabolism--ME; *Gene Products, tax--metabolism--ME; *HIV--physiology--PH; *HIV Long Terminal Repeat; *Human T-lymphotropic virus 1--genetics--GE; *Human T-lymphotropic virus 1--physiology--PH; *Regulatory Sequences, Nucleic Acid; *Repetitive Sequences, Nucleic Acid; *Virus Replication; Base Sequence; CD4-Positive T-Lymphocytes; Cell Line; Cell Survival; Chimera; Chloramphenicol O-Acetyltransferase--biosynthesis--BI; Gene Products, tat--biosynthesis--BI; Gene Products, tax--biosynthesis--BI; Gene Products, tax--genetics--GE; HIV--genetics--GE; Humans; Kinetics; Molecular Sequence Data; Plasmids; Recombinant Fusion Proteins--biosynthesis--BI; Research Support, Non-U.S. Gov't; Research Support, U.S. Gov't, P.H.S.; Restriction Mapping;

-more-

?

Display 6/9/2 (Item 1 from file: 154)

DIALOG(R) File 154:MEDLINE(R)

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Transfection

CAS Registry Number: 0 (Gene Products, tat); 0 (Gene Products, tax); 0 (Plasmids); 0 (Recombinant Fusion Proteins)

Enzyme Number: EC 2.3.1.28 (Chloramphenicol O-Acetyltransferase)

Record Date Created: 19951201

Record Date Completed: 19951201

- end of record -

? s s1 and pseudotyp?

296 S1

13362 PSEUDOTYP?

S7 19 S1 AND PSEUDOTYP?

? s s7 and tax

19 S7

90442 TAX

S8 0 S7 AND TAX

? s s1 and RNA (n) export?

296 S1

3941311 RNA

260647 EXPORT?

3371 RNA(N)EXPORT?

S9 4 S1 AND RNA (N) EXPORT?

? e au=harms, jerome

Ref	Items	Index-term
E1	2	AU=HARMS, JEANNE MCLAIN
E2	1	AU=HARMS, JEFFREY D.
E3	2	*AU=HARMS, JEROME
E4	4	AU=HARMS, JEROME S
E5	18	AU=HARMS, JEROME S.
E6	2	AU=HARMS, JEROME SCOTT
E7	2	AU=HARMS, JF
E8	1	AU=HARMS, JL
E9	1	AU=HARMS, JM
E10	5	AU=HARMS, JOACHIM
E11	1	AU=HARMS, JOAN YUKINO
E12	1	AU=HARMS, JOCHEN

Enter P or PAGE for more

? e au=harms jerome

Ref	Items	Index-term
E1	5	*AU=HARMS JEROME
E2	49	AU=HARMS JEROME S
E3	16	AU=HARMS JF
E4	3	AU=HARMS JH
E5	2	AU=HARMS JHK
E6	6	AU=HARMS JJ
E7	1	AU=HARMS JK
E8	5	AU=HARMS JL
E9	13	AU=HARMS JM
E10	2	AU=HARMS JOCHEN
E11	61	AU=HARMS JOERG
E12	23	AU=HARMS JOERG M

Enter P or PAGE for more

? e au=splitter, gary

Ref	Items	Index-term
E1	66	AU=SPLITTER, GA
E2	24	AU=SPLITTER, GA*
E3	12	*AU=SPLITTER, GARY
E4	24	AU=SPLITTER, GARY A
E5	88	AU=SPLITTER, GARY A.
E6	1	AU=SPLITTER, GARY ALLEN
E7	3	AU=SPLITTER, J. L.
E8	25	AU=SPLITTER, J. S.
E9	1	AU=SPLITTER, JACKIE LEE GOMER
E10	1	AU=SPLITTER, JANET L.
E11	1	AU=SPLITTER, JANET L. J.
E12	1	AU=SPLITTER, JANET LENA JANSSEN

Enter P or PAGE for more

? e au=splitter gary

Ref	Items	Index-term
E1	1	AU=SPLITTER G.S.
E2	163	AU=SPLITTER GA
E3	47	*AU=SPLITTER GARY
E4	178	AU=SPLITTER GARY A
E5	1	AU=SPLITTER GS
E6	2	AU=SPLITTER H
E7	1	AU=SPLITTER H W
E8	5	AU=SPLITTER J L
E9	4	AU=SPLITTER J S
E10	1	AU=SPLITTER J.L..
E11	2	AU=SPLITTER JL
E12	2	AU=SPLITTER JS

Enter P or PAGE for more

? e au=eakle, kurt

Ref	Items	Index-term
E1	6	*AU=EAKLE, KURT
E2	6	AU=EAKLE, KURT A
E3	15	AU=EAKLE, KURT A.
E4	4	AU=EAKLE, KURT ANDREW
E5	3	AU=EAKLE, MELISSA
E6	1	AU=EAKLE, N
E7	2	AU=EAKLE, N.
E8	2	AU=EAKLE, NORA
E9	4	AU=EAKLE, R. F.
E10	1	AU=EAKLE, RF

E11 1 AU=EAKLE, SUSAN D.
E12 1 AU=EAKLE, T. W

Enter P or PAGE for more
? e au=eakle kurt

Ref	Items	Index-term
E1	0	*AU=EAKLE KURT
E2	8	AU=EAKLE KURT A
E3	2	AU=EAKLE M.
E4	4	AU=EAKLE MELISSA
E5	3	AU=EAKLE N
E6	5	AU=EAKLE S
E7	4	AU=EAKLE S D
E8	1	AU=EAKLE S.
E9	2	AU=EAKLE S.D.
E10	2	AU=EAKLE SD
E11	4	AU=EAKLE STEPHAN
E12	6	AU=EAKLE T W

Enter P or PAGE for more
? e au=bremel, robert.

Ref	Items	Index-term
E1	18	AU=BREMEL, RD
E2	3	AU=BREMEL, RD*
E3	3	*AU=BREMEL, ROBERT
E4	10	AU=BREMEL, ROBERT D
E5	69	AU=BREMEL, ROBERT D.
E6	2	AU=BREMEL, ROBERT DUANE
E7	1	AU=BREMELL D
E8	2	AU=BREMELL D.
E9	5	AU=BREMELL DANIEL
E10	152	AU=BREMELL T
E11	56	AU=BREMELL T.
E12	2	AU=BREMELL THOMAS

Enter P or PAGE for more
? e au=bremel robert

Ref	Items	Index-term
E1	25	AU=BREMEL R.D.
E2	49	AU=BREMEL RD
E3	9	*AU=BREMEL ROBERT
E4	41	AU=BREMEL ROBERT D
E5	1	AU=BREMEL, D.
E6	15	AU=BREMEL, D. H.
E7	1	AU=BREMEL, D. R.
E8	1	AU=BREMEL, DAVID H
E9	2	AU=BREMEL, DAVID HERBERT
E10	1	AU=BREMEL, R
E11	87	AU=BREMEL, R. D.
E12	1	AU=BREMEL, R.-D.

Enter P or PAGE for more
?